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1645

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/833,675B

DATE: 02/05/2002
TIME: 14:04:35

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\02052002\I833675B.raw

ENTERED

2 <110> APPLICANT: Weimer, Thomas
 W--> 3 <120> TITLE OF INVENTION: PROCESS FOR FINDING OLIGONUCLEOTIDE SEQUENCES
 W--> 4 FOR NUCLEIC ACID AMPLIFICATION METHODS
 6 <130> FILE REFERENCE: 06478.1454-00
 W--> 7 <140> CURRENT APPLICATION NUMBER: 09/833,675B
 8 <141> CURRENT FILING DATE: 2001-04-13
 W--> 9 <160> NUMBER OF SEQ ID: 9
 10 <170> SOFTWARE: PatentIn version 3.1
 13 <210> SEQ ID NO: 1
 14 <211> LENGTH: 27
 15 <212> TYPE: DNA
 16 <213> ORGANISM: Red Sea Bream Iridovirus
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 19 <222> LOCATION: (1)..(27)
 20 <223> OTHER INFORMATION: n=inosine
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 28 <213> ORGANISM: Homo sapiens
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 30 actccaccat agatcact 18
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 39 <223> OTHER INFORMATION: n=inosine
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 45 <211> LENGTH: 21
 46 <212> TYPE: DNA
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 50 <222> LOCATION: (1)..(21)
 51 <223> OTHER INFORMATION: n=inosine
 W--> 52 <400> SEQUENCE: 4

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Input Set : A:\PT0.VSK.txt
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 57 <211> LENGTH: 20
 58 <212> TYPE: DNA
 59 <213> ORGANISM: Helianthus tuberosus
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 62 <222> LOCATION: (1)..(20)
 63 <223> OTHER INFORMATION: n=inosine
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 69 <211> LENGTH: 25
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 75 <223> OTHER INFORMATION: n=inosine
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 109 <222> LOCATION: (1)..(31)
 110 <223> OTHER INFORMATION: n=inosine

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Input Set : A:\PTO.VSK.txt
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W¹ 111 <400> SEQUENCE: 9
W² 112 gtacctgggt cncgaaagnc cttgtggta c t

31

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/833,675B

DATE: 02/05/2002

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Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\02052002\I833675B.raw

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L:7 M:283 W: Missing Blank Line separator, <140> field identifier
L:9 M:283 W: Missing Blank Line separator, <160> field identifier
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L:22 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1
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L:41 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
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L:53 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4
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L:65 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5
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L:77 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
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L:89 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7
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L:111 M:283 W: Missing Blank Line separator, <400> field identifier
L:112 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9